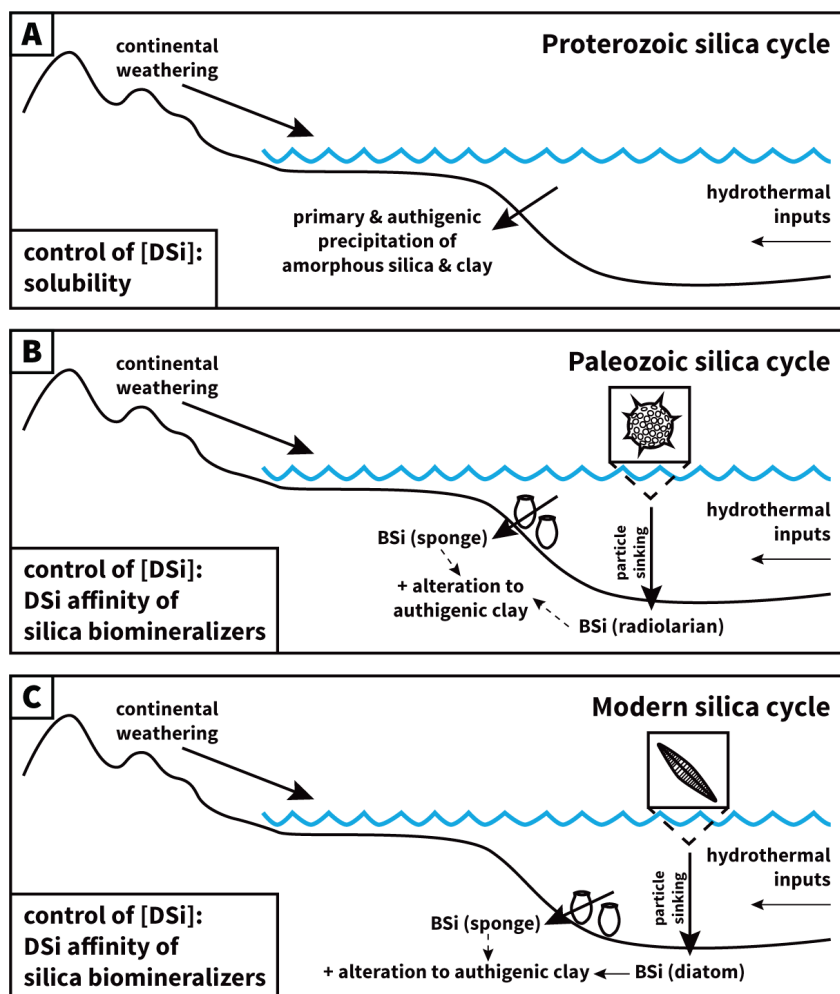


## Supplementary Information for:

### Isotopic analyses of Ordovician-Silurian siliceous skeletons indicate silica-depleted Paleozoic oceans

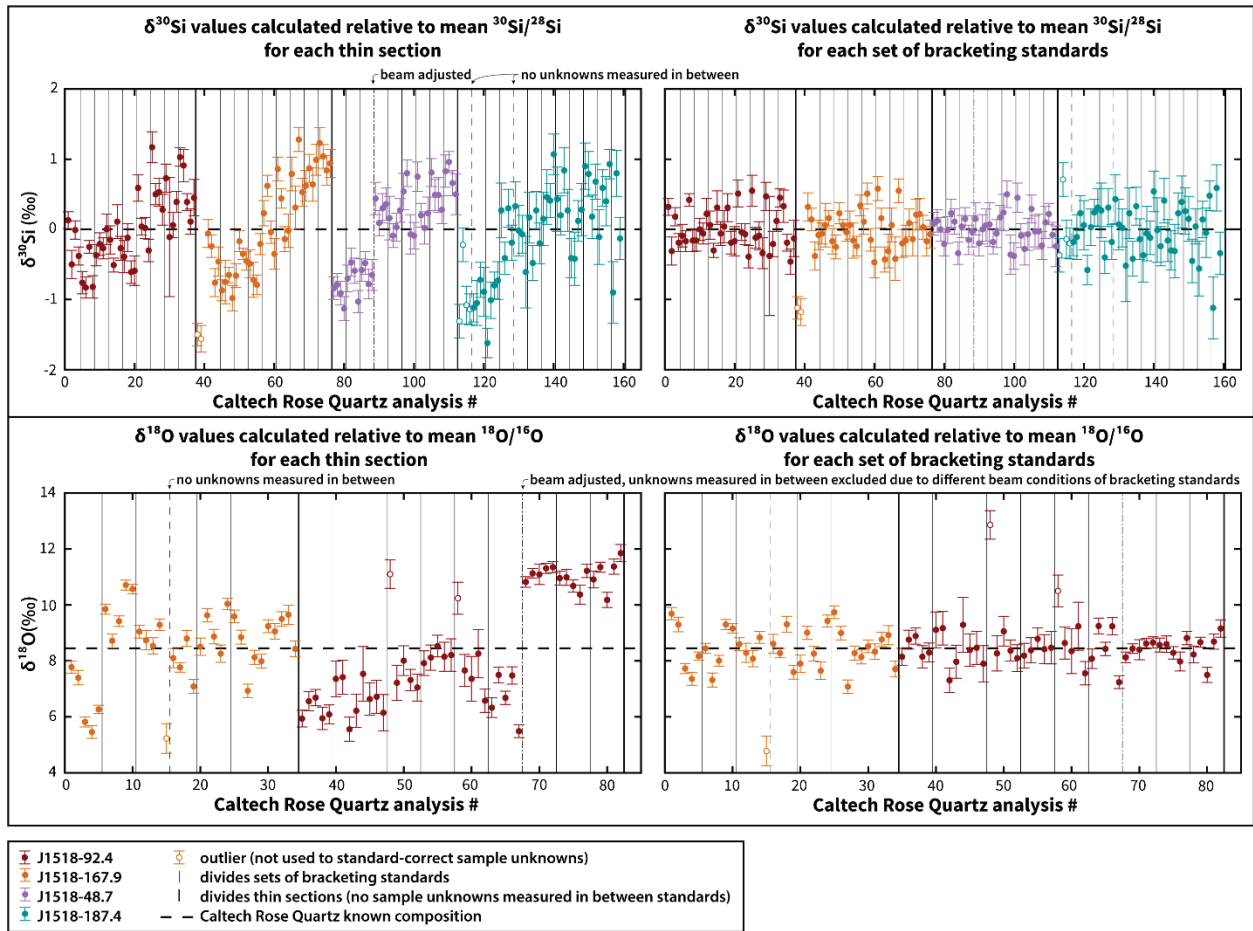
#### Contents

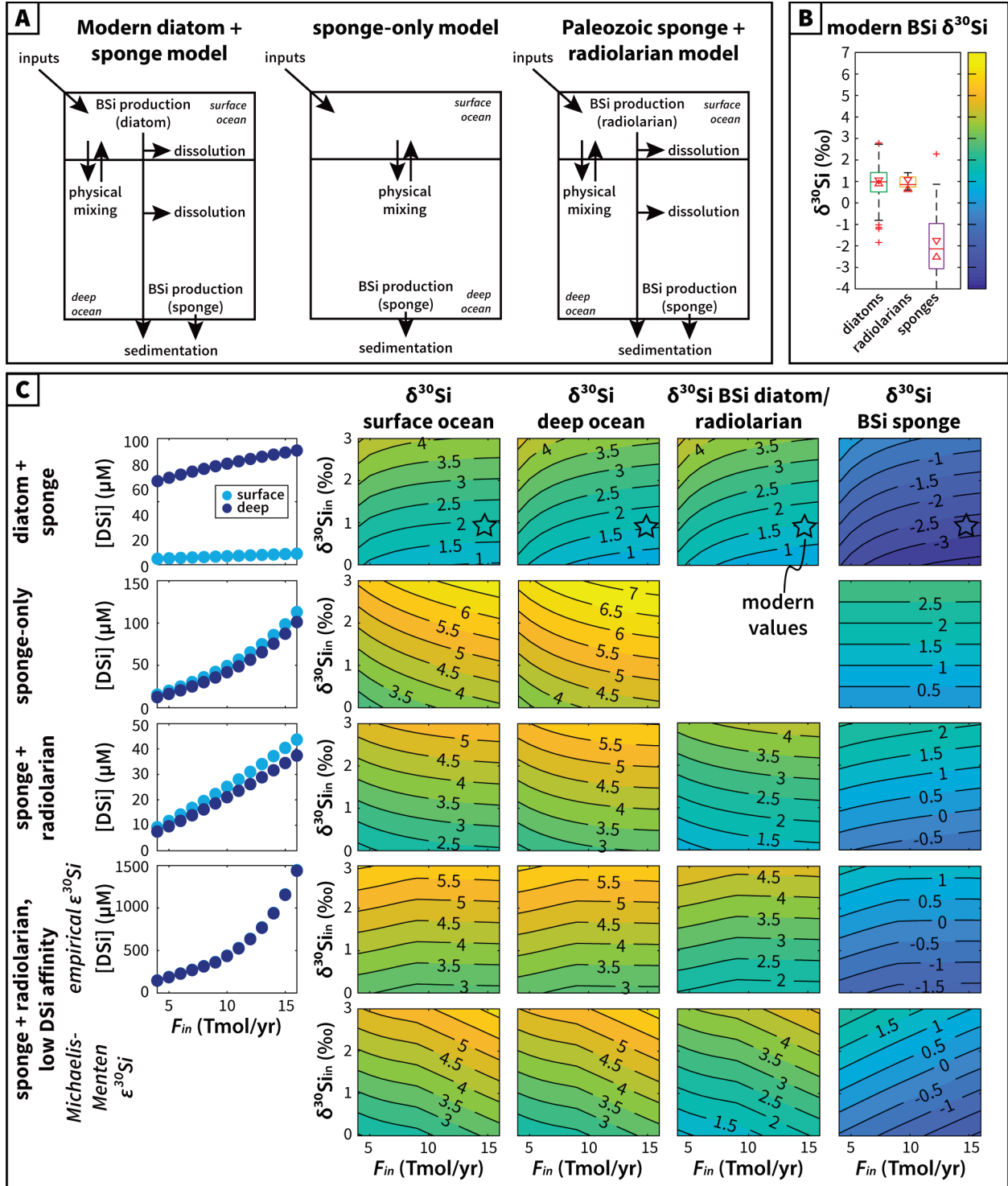
This supplementary information document includes 12 figures and 2 tables. Dataset S1 is provided in a separate .xlsx file.



**Figure S1**—Simplified schematic illustrations of the Proterozoic (A), Paleozoic (B), and Modern (C) marine silica cycle. (A) In Proterozoic oceans, [DSi] is thought to have been controlled primarily by mineral solubilities—both amorphous silica and authigenic clay minerals—which were the major sinks for DSi delivered to seawater from continental weathering and hydrothermal inputs (Siever 1992). (B) In Paleozoic oceans, the marine silica cycle was impacted by the evolution of silica-biomineralizing organisms. BSi production in the water column (radiolarians) and on the seafloor (sponges) added new major sinks to balance DSi inputs. Early diagenetic alteration of these silica biominerals to authigenic clay minerals via reverse weathering reactions could have constituted an additional silica sink, but the reactivity of sponge spicules and radiolarian tests has not been well studied. (C) In modern oceans, BSi production in the water column (diatoms) and on the seafloor (sponges) constitute major silica sinks; precipitation of authigenic clay minerals via alteration of diatom frustules (i.e., reverse weathering) is an

additional important sink in the modern marine silica cycle (Rahman, Aller, and Cochran 2017; Tréguer and De La Rocha 2013).

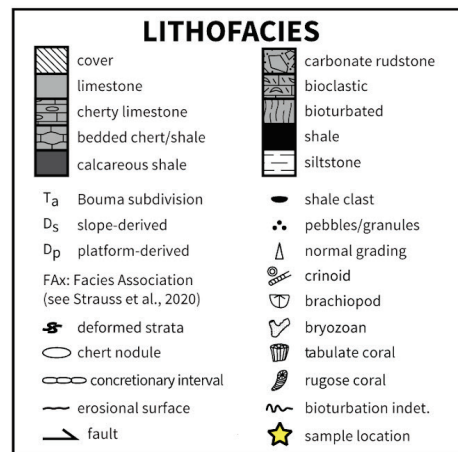
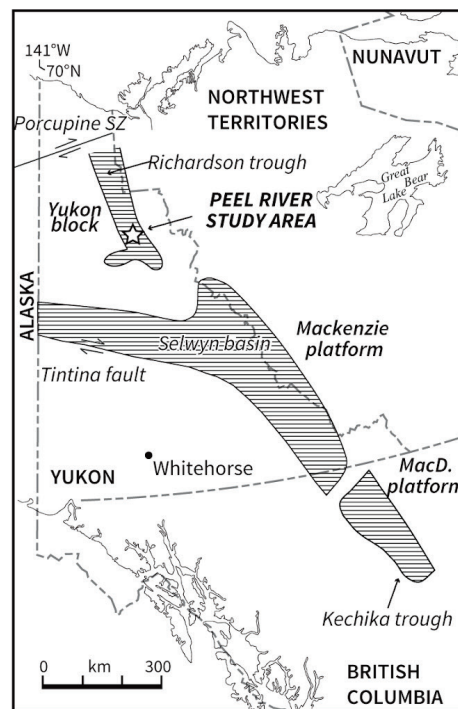
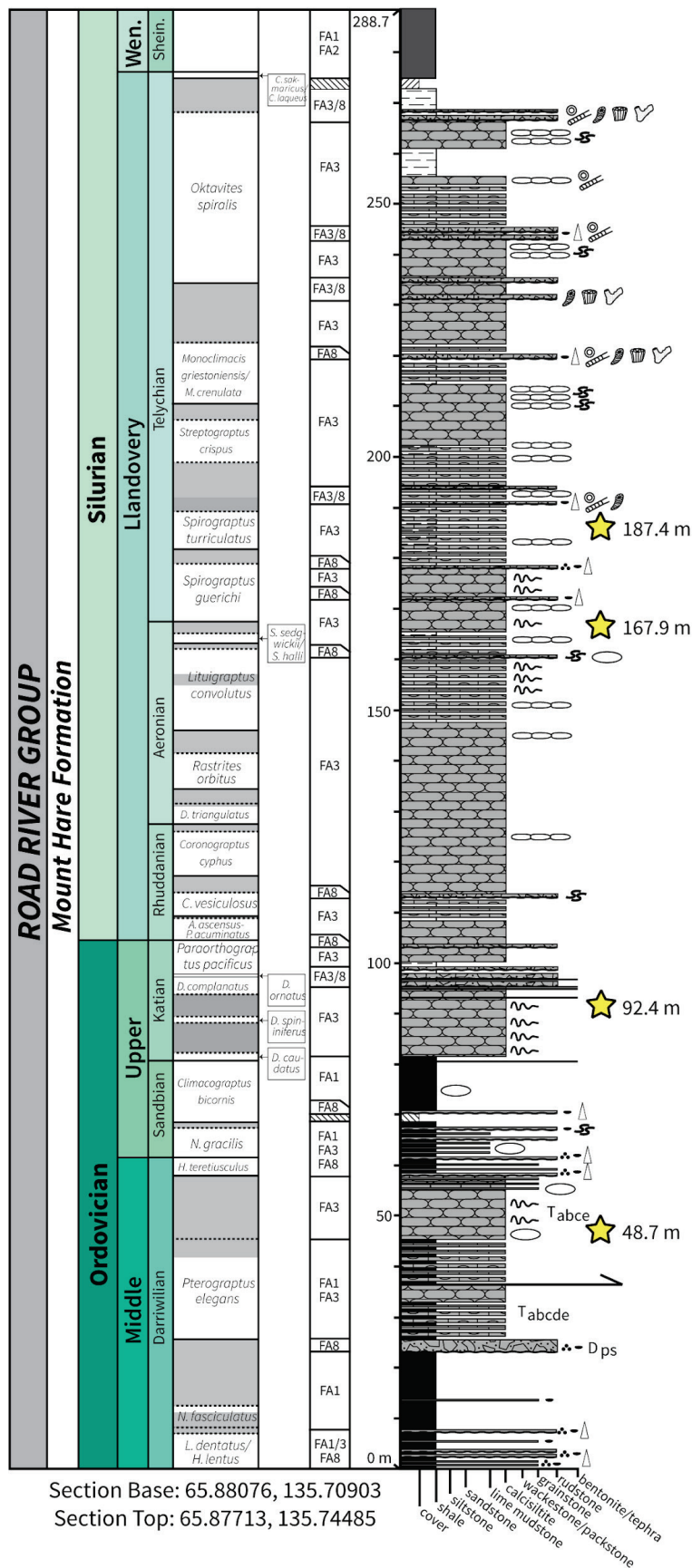




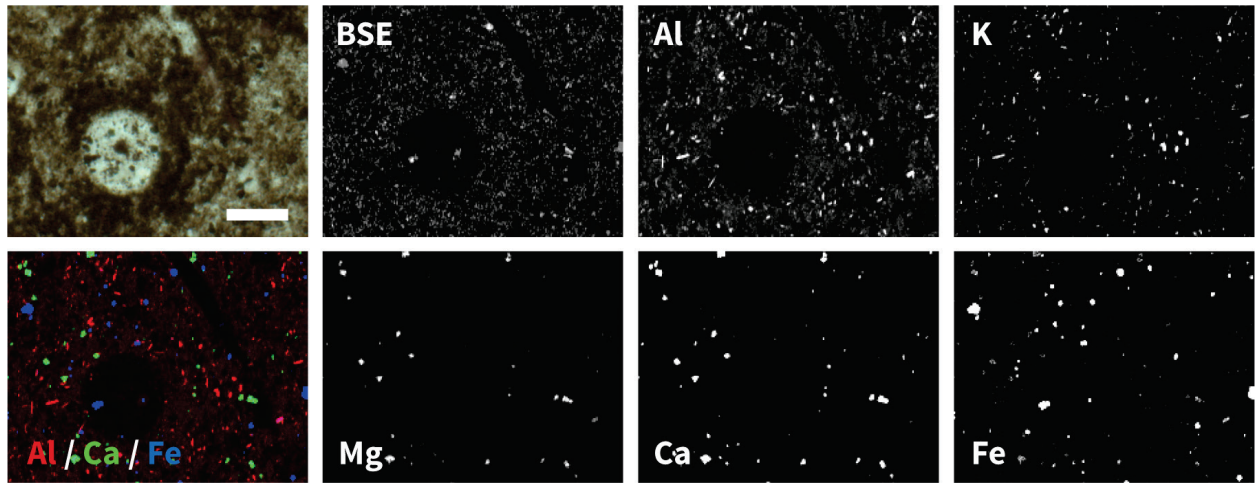
**Figure S3**—(A) Schematic illustrations of marine silica cycle models. (B) Box plots of  $\delta^{30}\text{Si}$  distributions in modern marine BSi—diatoms (Cardinal et al. 2007; Fripiat et al. 2011; Hendry and Robinson 2012; De La Rocha, Brzezinski, and DeNiro 2000; Egan et al. 2012; Varela, Pride, and Brzezinski 2004), radiolarians (Abelmann et al. 2015), and sponge spicules (De La Rocha 2003; Hendry and Robinson 2012; Wille et al. 2010; Douthitt 1982) plotted next to the colorbar scale shown in (C). (C) Box model predictions: scatter plots of  $[\text{DSi}]$  for surface and deep oceans and contour plots  $\delta^{30}\text{Si}$  values of surface

ocean, deep ocean, planktonic silica biomineralizers (diatoms or radiolarians), and sponges for five versions of the box model, from top to bottom: (1) diatom + sponge (with stars illustrating corresponding modern  $\delta^{30}\text{Si}$  values colored following the same scale); (2) sponge-only; (3) sponge + radiolarian assuming modern DSi affinity and modern empirical  $\epsilon^{30}\text{Si}$ -[DSi] calibration; (4) sponge + radiolarian assuming 20x lower DSi affinity than modern sponges and modern  $\epsilon^{30}\text{Si}$ -[DSi] calibration; and (5) same as (4) but with Michaelis-Menten  $\epsilon^{30}\text{Si}$ -[DSi] prediction. All plots share the same color scale shown in (B). [DSi] predictions are not shown for scenario (5) because they are identical to scenario (4).



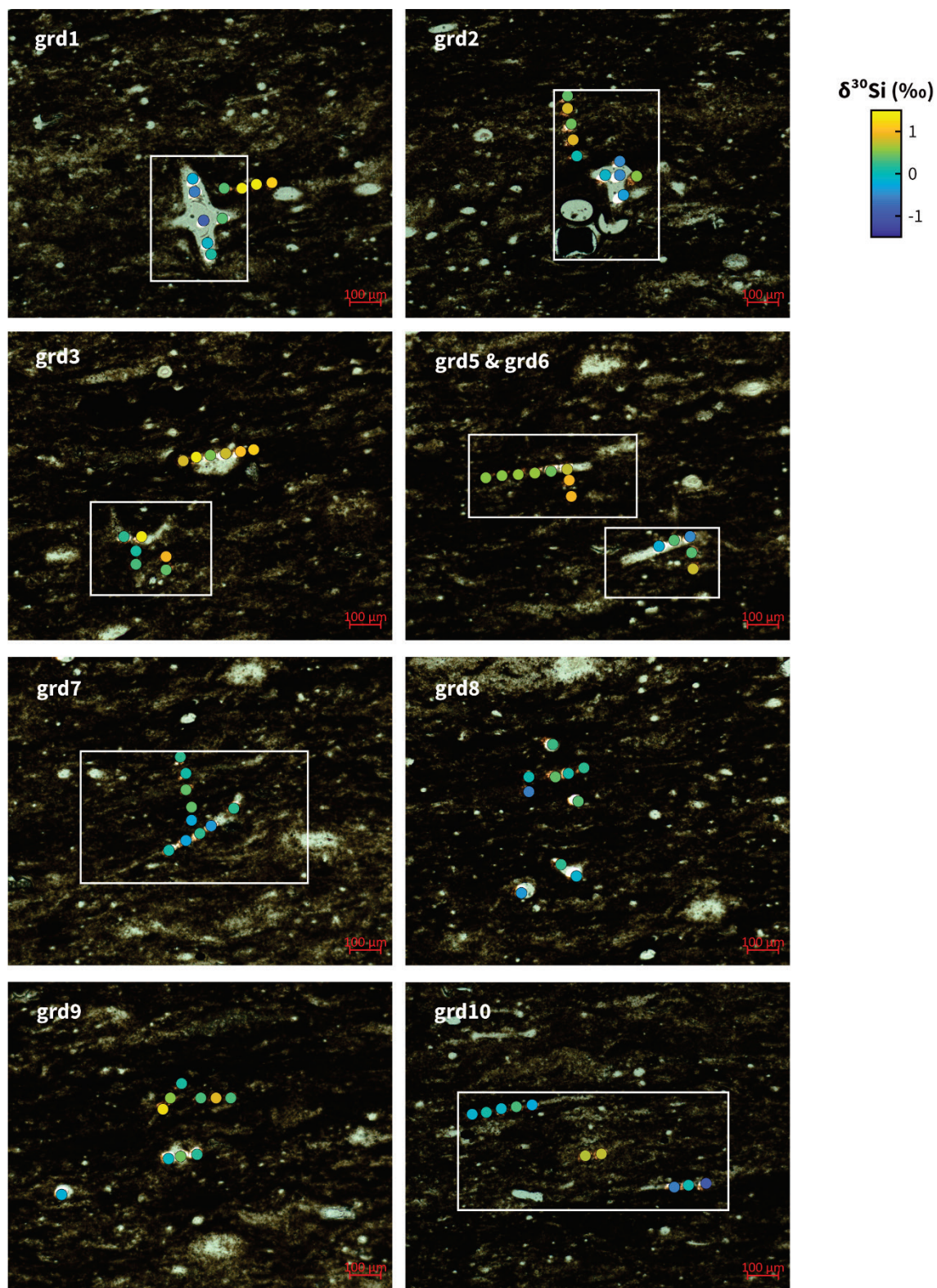


**Figure S4**—Location map of study area and stratigraphic column of section J1518 from Strauss et al. (2020) illustrating stratigraphic heights, sedimentary facies, and ages of samples in this study.



**Figure S5**—Additional electron microprobe elemental data of radiolarian and spicule from sample J1518-92.4, comparing transmitted light image (upper left); back-scattered electron (BSE) image; maps of elements Al, K, Mg, Ca, and Fe; and an RGB map plotting Al (red), Ca (green), and Fe (blue). Scale bar is 100  $\mu\text{m}$ .

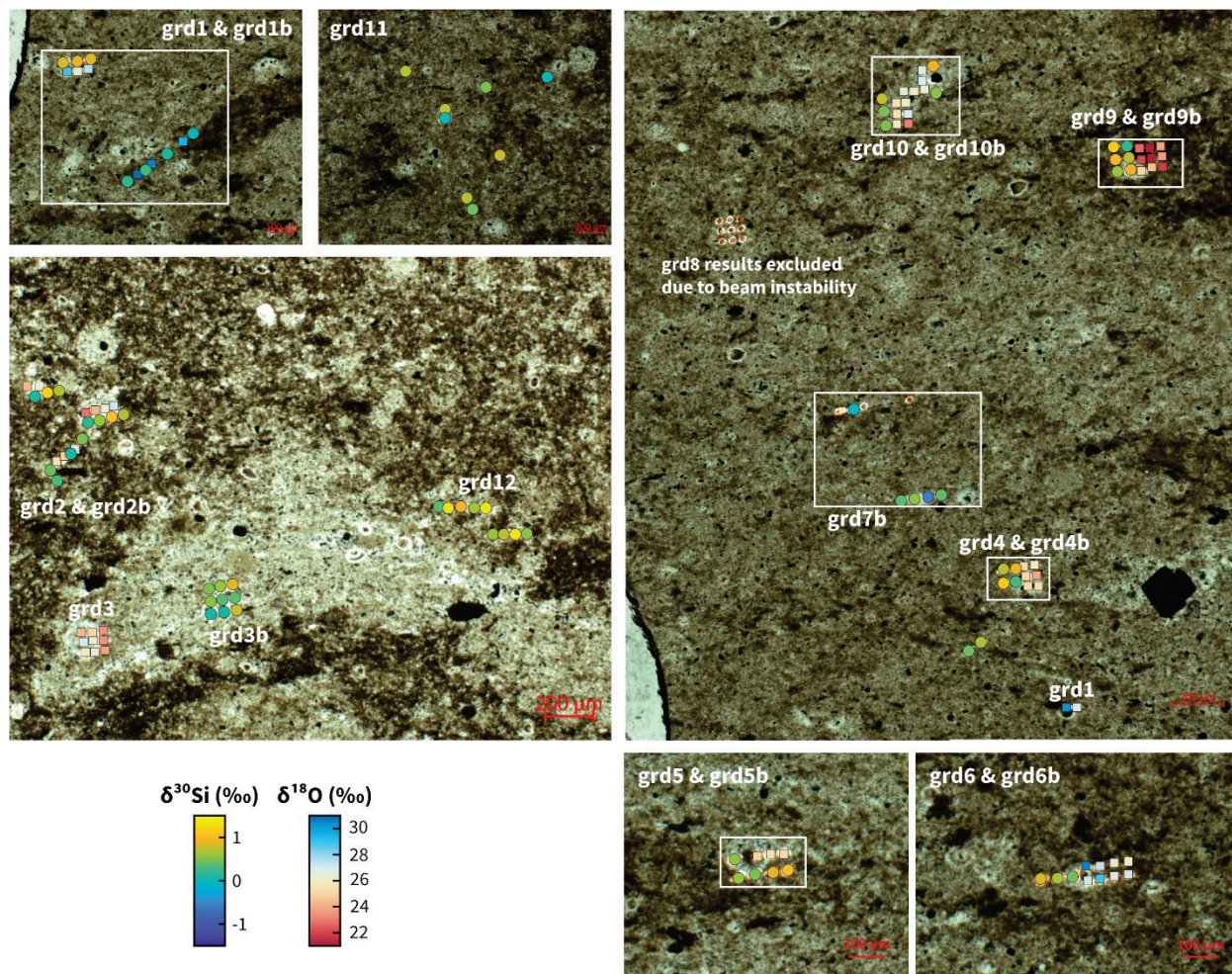




**Figure S6**—Plane-polarized transmitted light images of sample J1518 48.7 showing locations of  $\delta^{30}\text{Si}$  analytical spots (circles, color-mapped by  $\delta^{30}\text{Si}$  value). White boxes indicate approximate boundaries of EMP WDS maps collected from this sample. A higher resolution version of this image, a high resolution

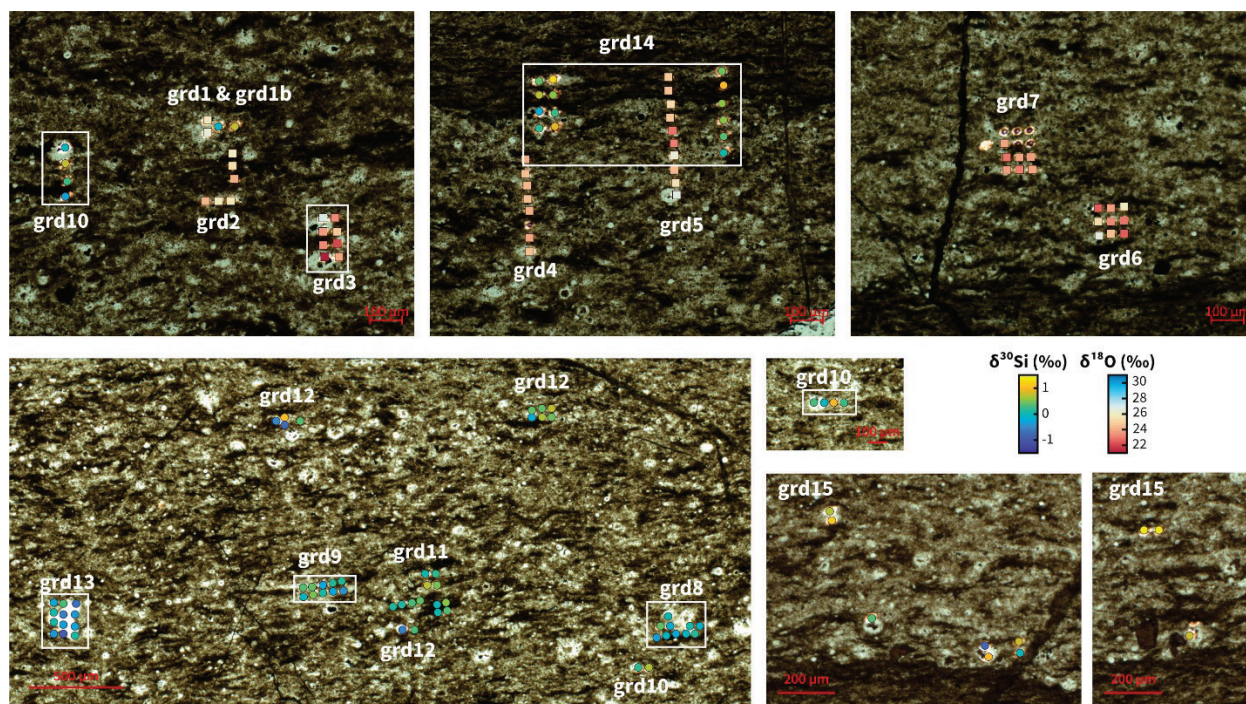


map of the entire sample, and EMP WDS maps are archived in the linked OSF data repository at doi: 10.17605/OSF.IO/4AES6.



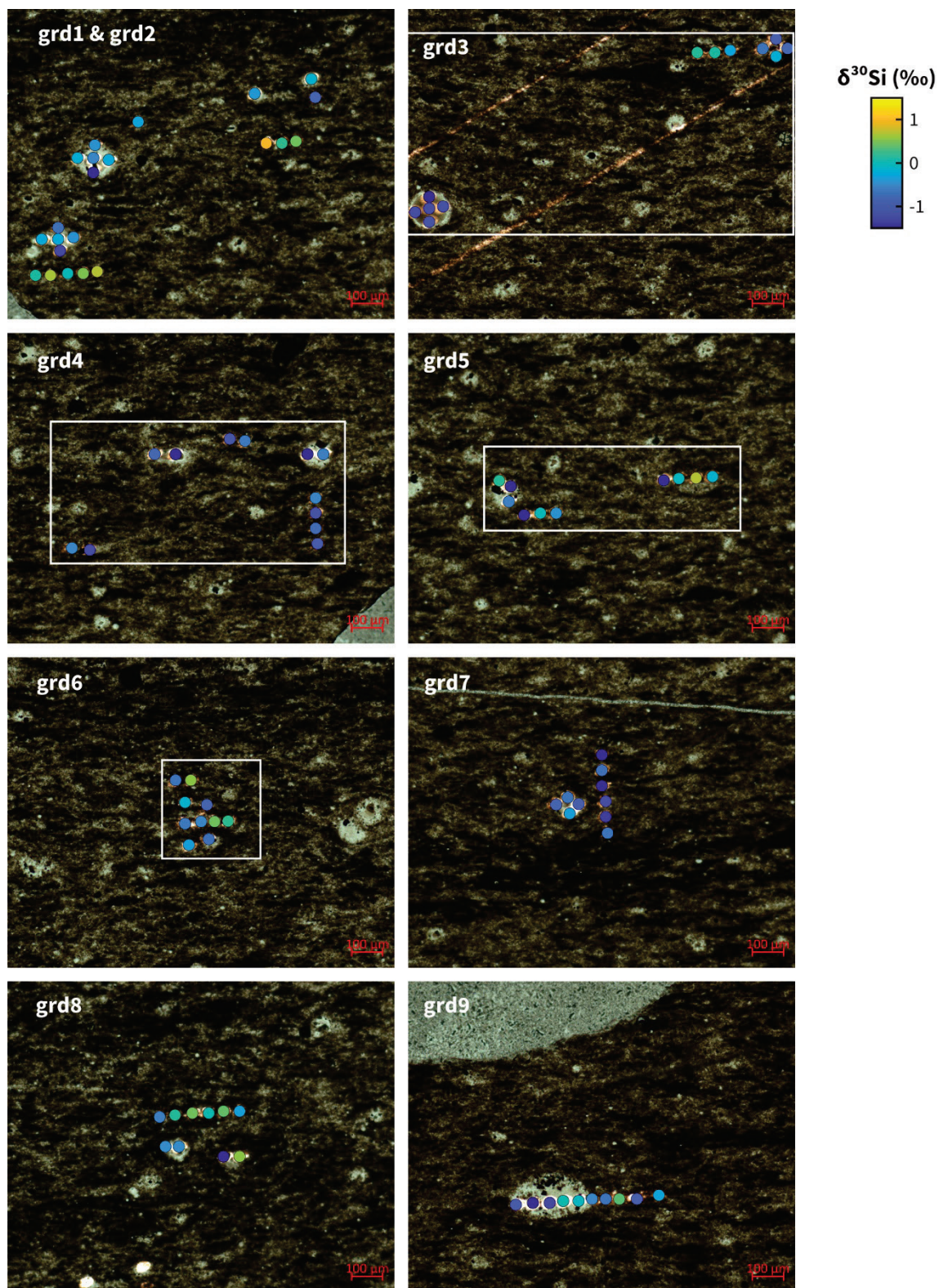
**Figure S7**— Plane-polarized transmitted light images of sample J1518 92.4 showing locations of  $\delta^{30}\text{Si}$  and  $\delta^{18}\text{O}$  analytical spots ( $\delta^{30}\text{Si}$ : circles, color-mapped by  $\delta^{30}\text{Si}$  value;  $\delta^{18}\text{O}$ : squares, color-mapped by  $\delta^{18}\text{O}$  value). White boxes indicate approximate boundaries of EMP WDS maps collected from this sample. A higher resolution version of this image, a high resolution map of the entire sample, and EMP WDS maps are archived in the linked OSF data repository at doi: 10.17605/OSF.IO/4AES6.





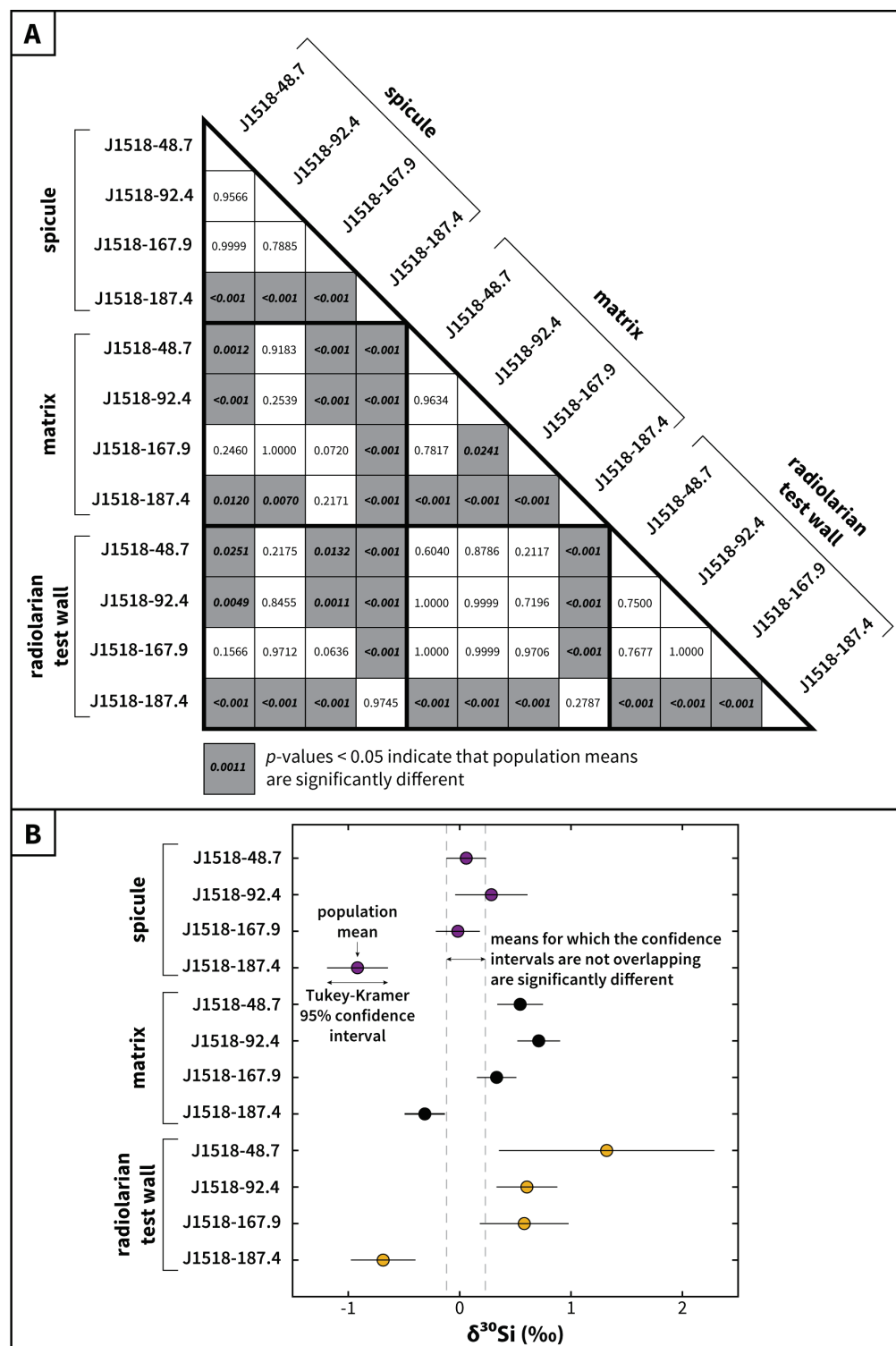
**Figure S8**— Plane-polarized transmitted light images of sample J1518 167.9 showing locations of  $\delta^{30}\text{Si}$  and  $\delta^{18}\text{O}$  analytical spots ( $\delta^{30}\text{Si}$ : circles, color-mapped by  $\delta^{30}\text{Si}$  value;  $\delta^{18}\text{O}$ : squares, color-mapped by  $\delta^{18}\text{O}$  value). White boxes indicate approximate boundaries of EMP WDS maps collected from this sample. A higher resolution version of this image, a high resolution map of the entire sample, and EMP WDS maps are archived in the linked OSF data repository at doi: 10.17605/OSF.IO/4AES6.





**Figure S9**— Plane-polarized transmitted light images of sample J1518 187.4 showing locations of  $\delta^{30}\text{Si}$  (circles, color-mapped by  $\delta^{30}\text{Si}$  value). White boxes indicate approximate boundaries of EMP WDS maps collected from this sample. A higher resolution version of this image, a high resolution map of the entire

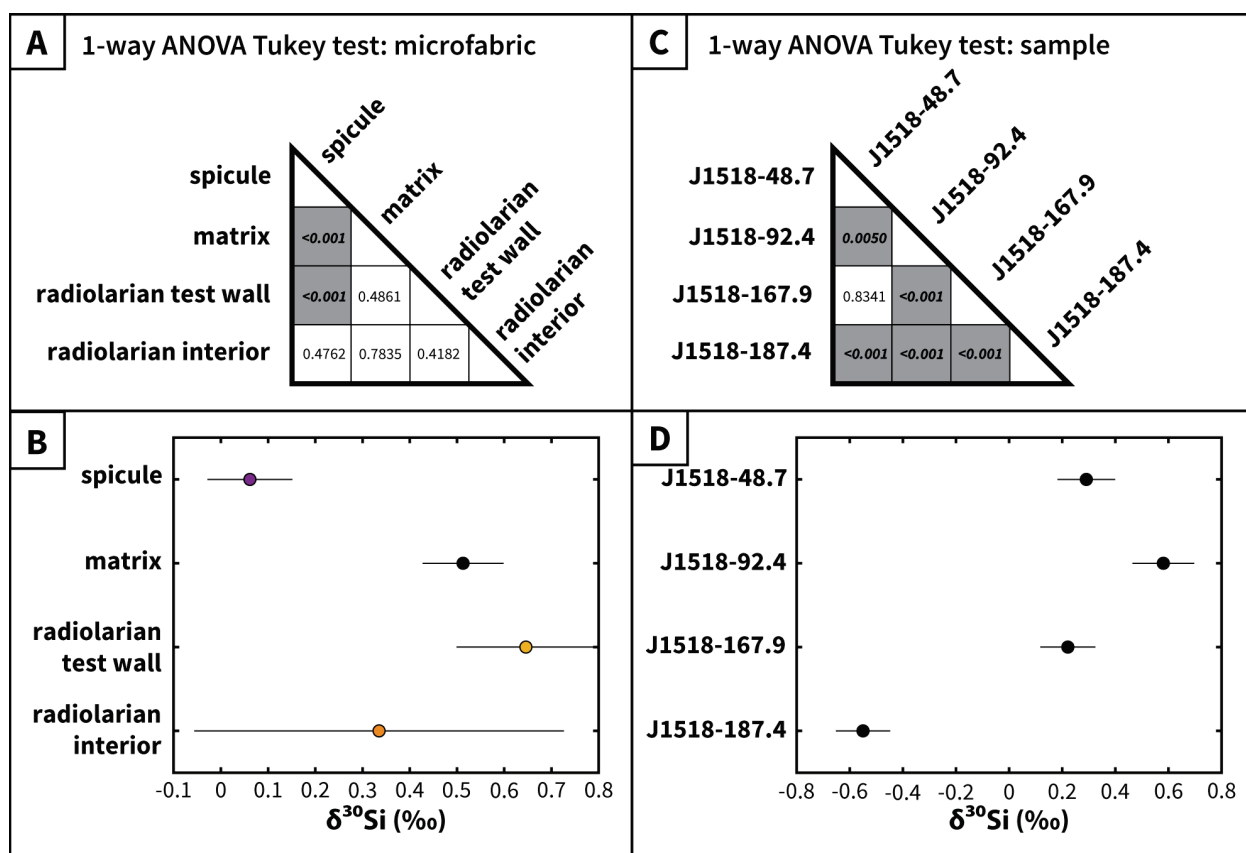
sample, and EMP WDS maps are archived in the linked OSF data repository at doi: 10.17605/OSF.IO/4AES6.



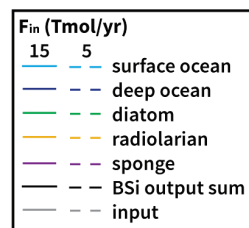
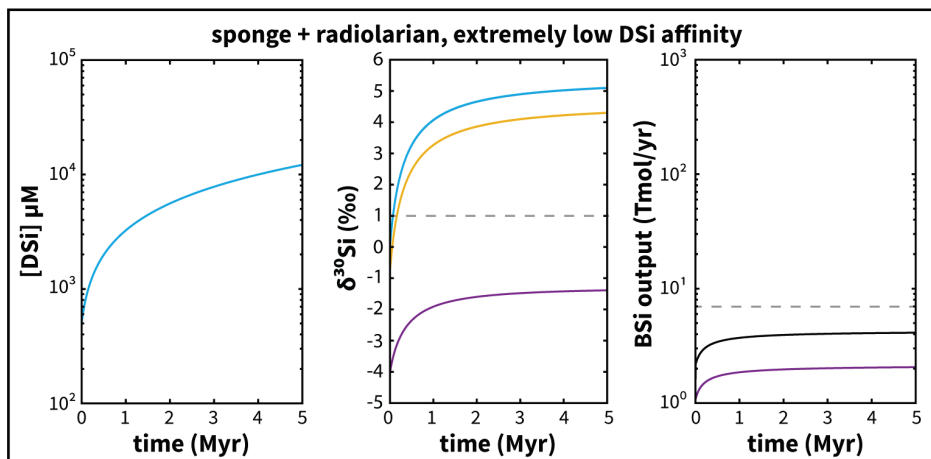
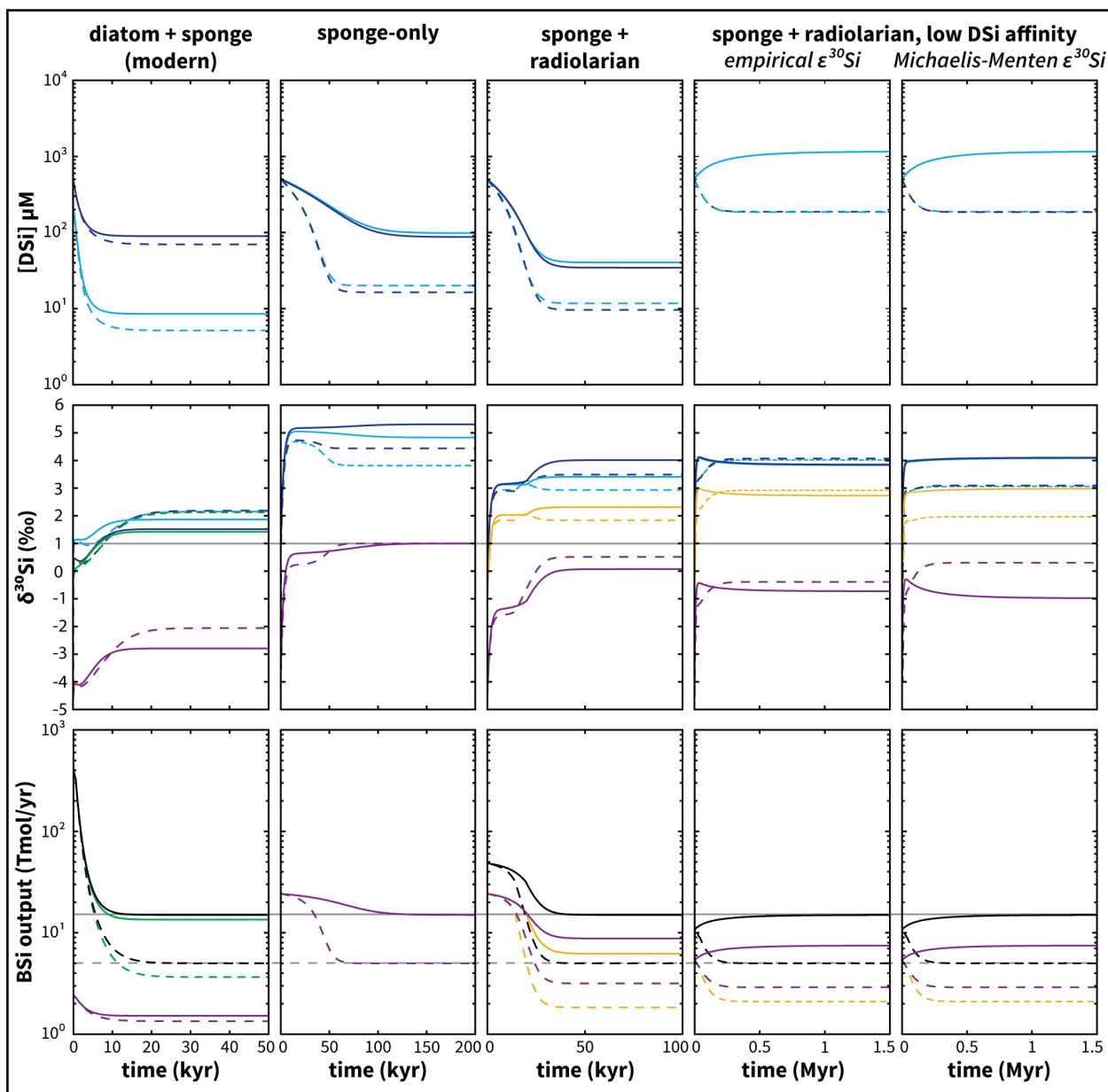
**Figure S10**— Results of post-hoc multiple comparison analysis (following the Tukey-Kramer procedure) based on an unbalanced 2-way ANOVA (ANalysis Of Variance) performed on SIMS  $\delta^{30}\text{Si}$  data grouped

by sample ID and microfabric (spicule, matrix, radiolarian test wall; radiolarian interiors were excluded because no spots of this type were analyzed in sample J1518-167.9). (A) Matrix of  $p$ -values for every pair of sample-microfabric combination; grey cells with bold italic text indicate combinations for which the population means are significantly different. (B) Alternative visualization of the same information as in (A), where colored circles indicate population means and horizontal lines indicate 95% confidence intervals; means for which the confidence intervals do not overlap are significantly different (e.g., means for which the confidence intervals are outside the dashed grey lines are significantly different from spicules in sample J1518-48.7). This analysis indicated that the populations of spicules within samples J1518-48.7, J1518-92.4, and J1518-167.9 were not significantly different from each other and that the populations of radiolarian test walls within samples J1518-48.7, J1518-92.4, and J1518-167.9 were not significantly different from each other. For this ANOVA, the F-values were 70.59, 28.37, and 2.34 (corresponding  $p$ -values  $4.9 \times 10^{-35}$ ,  $4.6 \times 10^{-12}$ , and 0.0315) for groupings of sample ID, microfabric, and sample ID \* microfabric, respectively—indicating that differences between means grouped by sample, microfabric, and microfabric within sample are significant.





**Figure S11**— Results of post-hoc multiple comparison analysis (following the Tukey-Kramer procedure) based on an unbalanced 1-way ANOVA on SIMS  $\delta^{30}\text{Si}$  data grouped by microfabric (A, B) and sample (C, D), following the same figure design as in Figure S10 and described in the Figure S10 caption. (A, B) Here, data were grouped only by microfabric, so analyses of the same microfabric are combined across all samples. Sample J1518-187.4 was excluded from this analysis because the Tukey test of the unbalanced 2-way ANOVA (Figure S10) demonstrated that the means of all microfabrics in this sample were significantly different from the means of all microfabrics in the remaining samples (except for the pair J1518-187.4 matrix – J1518-167.9 spicule), supporting the interpretation that this sample may have been more strongly affected by diagenesis. This analysis indicated that the population means of spicules and radiolarian test walls among these three samples are significantly different. For this ANOVA, the F-value was 22.46 and the  $p$ -value was  $6.7 \times 10^{-13}$ , indicating that differences between means grouped by microfabric were significant. (C, D) Here, data were grouped only by sample, so analyses of all microfabrics in each sample were combined. This analysis indicated that the population means (of all microfabric types) were significantly different among all pairs of samples except for the pair J1518-48.7 – J1518-167.9. We attributed some of these differences to the different numbers of analyses of each fabric type within each sample (i.e., different mixtures of the apparent end-member fabric compositions illustrated in panel B) and apparent diagenetic influence on sample J1518-187.4. For this ANOVA, the F-value was 68.43 and the  $p$ -value was  $9.999 \times 10^{-39}$ , indicating that differences between means grouped by sample were significant.



**Figure S12**—Evolution of [DSi],  $\delta^{30}\text{Si}$ , and BSi output flux over time for different box model versions, illustrating approach to steady state. The plots in the upper panel correspond to the models shown in Figure S3A for  $F_{in} = 5$  Tmol/yr (dashed lines) or 15 Tmol/yr (solid lines) and  $\delta^{30}\text{Si}_{in} = 1\text{‰}$ . The plots in the lower panel illustrate similar model parameters to those in the low DSi affinity with Michaelis-Menten  $\epsilon^{30}\text{Si}$  from the upper panel, but with DSi affinity reduced further. Under these conditions, [DSi] can exceed 1000  $\mu\text{M}$ , but the model fails to reach steady state after 5 Ma (a significantly longer duration than in the models in the upper panel) because BSi production cannot keep pace with  $F_{in}$  and [DSi] continues to rise beyond abiotic mineral saturation thresholds (abiotic precipitation is not included in this model). The plots in the upper panel also illustrate that  $\delta^{30}\text{Si}_{\text{sponge}}$  only transiently records extremely low values  $\sim -4\text{‰}$  and that under Paleozoic conditions, we expect  $\delta^{30}\text{Si}_{\text{sponge}} > -2\text{‰}$  and that both surface and deep ocean  $\delta^{30}\text{Si}_{\text{DSi}}$  were  $\geq 3\text{‰}$ .

**Table S1**—Parameters for box models

| Parameter   | Diatom + sponge model   | Sponge-only model  | Sponge + radiolarian modern DSi affinity model  | Sponge + radiolarian low DSi affinity model  |
|---|---|--|---|--|
| Ocean volume ( $\text{m}^3$ )                                     | $1.37 \times 10^{18}$   |  |   |  |
| Ocean surface area ( $\text{m}^2$ )                               | $3.6 \times 10^4$   |  |   |  |
| Fraction of euphotic zone (dimensionless)                         | 0.0263  |  |   |  |
| Physical mixing flux ( $\text{m}^3/\text{yr}$ )                   | $1.37 \times 10^{18}$ (De La Rocha and Bickle 2005)   |  |   |  |
| Particle sinking velocity, surface ocean ( $\text{m}/\text{yr}$ ) | 1800 (Passow, French, and Robert 2011)  |  |   |  |
| Particle sinking velocity, deep ocean ( $\text{m}/\text{yr}$ )    | 73000 (Passow, French, and Robert 2011)   |  |   |  |
| Maximum BSi production rate, $V_{max}$                            | Diatoms: $5 \times 10^{14}$ mol/yr (Sarmiento and Gruber 2006); sponges: 0.13 $\mu\text{mol}/\text{hr}/\text{mL}$ sponge volume (Maldonado et al. 2011) | 0.13 $\mu\text{mol}/\text{hr}/\text{mL}$ sponge volume (Maldonado et al. 2011) | 0.13 $\mu\text{mol}/\text{hr}/\text{mL}$ sponge volume (Maldonado et al. 2011); radiolarians assumed to have same $V_{max}$ when scaled to mol/yr | 0.05 $\mu\text{mol}/\text{hr}/\text{mL}$ sponge volume; $\sim 2.5\times$ lower than the lowest value observed in modern sponges (López-Acosta et al. 2018) |

|  |  |   |   |  |
|--|--|---|---|--|
| Sponge seafloor density  | 0.34 L sponge / m <sup>2</sup> ; sponge-poor oligotrophic Mediterranean estimate (Maldonado et al. 2011) |   |   |  |
| Fraction of seafloor colonized by sponges                          | 0.025  | 0.2   |   |  |
| $K_m$ (μM)   | Diatom: 9 (Nelson et al. 1995; Amo and Brzezinski 1999); sponge: 75 (Maldonado et al. 2011)              | 75 (Maldonado et al. 2011)  | 75 (Maldonado et al. 2011); radiolarians assumed to have the same $K_m$   | 500; ~5x higher than the highest value observed in modern sponges (López-Acosta et al. 2018)   |
| BSi dissolution rate constant, surface ocean, $k_{surface}$ (1/yr) | 9 (Frings et al. 2016) (applies only to radiolarians)  | N/A   | 9 (applies only to radiolarians)  |  |
| BSi dissolution rate constant, surface ocean, $k_{deep}$ (1/yr)    | 24 (Frings et al. 2016) (applies only to radiolarians)   | N/A   | 3 (applies only to radiolarians) (Erez, Takahashi, and Honjo 1982; Berger 1968)                                 |  |
| Apparent BSi solubility, $C_{eq}$ (μM)                             | 350 (Loucaides et al. 2012)  | N/A   | 100 (applies only to radiolarians) (Berger 1968)  | 350 (applies only to radiolarians)   |
| $\epsilon^{30}\text{Si}$ (‰)                                       | diatom: -1.1 (de la Rocha, Brzezinski, and DeNiro 1997; Sutton et al. 2013; Milligan et al. 2004)        | sponge: $-4.6 + \frac{27.6}{7.4 + [\text{DSi}]}$ (Hendry et al. 2019) | sponge: $-4.6 + \frac{27.6}{7.4 + [\text{DSi}]}$ (Hendry et al. 2019); radiolarian: -1.1 (Abelmann et al. 2015) | sponge: $-4.6 + \frac{27.6}{7.4 + [\text{DSi}]}$ (Hendry et al. 2019) or extrapolation of Michaelis-Menten relationship for high [DSi]* (Wille et al. 2010; Cassarino et al. 2018); radiolarian: -1.1 (Abelmann et al. 2015) |

\*Following Wille et al. (2010) and Cassarino et al. (2018), we solve for  $\epsilon^{30}\text{Si} = \epsilon_{tl} + (\epsilon_p -$

$$\epsilon_E) \left( I - \frac{\frac{V_{max,P}}{(\frac{K_{m,P}}{[\text{DSi}]} + 1)}}{\frac{V_{max,I}}{(\frac{K_{m,I}}{[\text{DSi}]} + 1)}} \right), \text{ where } \epsilon_{tl} = -1.34\text{‰ is the fractionation during DSi uptake, } (\epsilon_p - \epsilon_E) = -5.39\text{‰ is the}$$

difference between fractionation during polymerization ( $\epsilon_p$ ) and fractionation during efflux ( $\epsilon_E$ ),  $V_{max,P}$  is the maximum polymerization rate (set as the maximum BSi production rate given in the table),  $V_{max,I}$  is

the maximum incorporation rate estimated as 9  $\mu\text{mol/hr/mL}$  sponge following Cassarino et al. (2018),  $K_{m,P}$  is the half saturation constant of polymerization (set as the  $K_m$  value in the table), and  $K_{m,I}$  is the half saturation constant of incorporation, calculated as  $\frac{V_{max,I}K_{m,P}}{V_{max,P}}$ .

**Table S2—SIMS  $\delta^{30}\text{Si}$  and  $\delta^{18}\text{O}$  overview statistics**

|   | J1518-48.7 | J1518-92.4 | J1518-167.9 | J1518-187.4 |
|---|------------|------------|-------------|-------------|
| <b><math>\delta^{30}\text{Si}</math> analyses</b> |            |            |             |             |
| <i>radiolarian interior</i>                       |            |            |             |             |
| n analyses  | 2          | 4          | 0           | 11          |
| median (‰)  | 0.65       | 0.25       | N/A         | -0.98       |
| standard deviation (‰)                            | 0.19       | 0.59       | N/A         | 0.61        |
| maximum (‰)                                       | 0.78       | 0.80       | N/A         | 0.01        |
| minimum (‰)                                       | 0.51       | -0.57      | N/A         | -2.15       |
| <i>radiolarian test wall</i>                      |            |            |             |             |
| n analyses  | 2          | 18         | 9           | 16          |
| median (‰)  | 1.32       | 0.57       | 0.70        | -0.64       |
| standard deviation (‰)                            | 0.38       | 0.24       | 0.47        | 0.52        |
| maximum (‰)                                       | 1.59       | 1.06       | 1.15        | -0.05       |
| minimum (‰)                                       | 1.05       | 0.13       | -0.11       | -1.88       |
| <i>spicule</i>                                    |            |            |             |             |
| n analyses  | 48         | 13         | 36          | 18          |
| median (‰)  | 0.09       | 0.24       | 0.05        | -1.03       |
| standard deviation (‰)                            | 0.49       | 0.26       | 0.37        | 0.71        |
| maximum (‰)                                       | 1.51       | 0.70       | 0.82        | 0.52        |
| minimum (‰)                                       | -0.98      | -0.11      | -0.98       | -2.40       |

|   |       |       |       |       |
|---|-------|-------|-------|-------|
| <i>matrix</i>   |       |       |       |       |
| n analyses  | 33    | 39    | 48    | 51    |
| median (‰)  | 0.41  | 0.67  | 0.52  | -0.37 |
| standard deviation (‰)                                  | 0.47  | 0.35  | 0.47  | 0.65  |
| maximum (‰)   | 1.54  | 1.54  | 1.60  | 0.67  |
| minimum (‰)   | -0.59 | 0.09  | 0.80  | -2.10 |
| $\Delta^{30}\text{Si}_{\text{sponge-radiolarian}}$ (‰)  |       |       |       |       |
| w/ radiolarian test wall median $\delta^{30}\text{Si}$  | -1.23 | -0.33 | -0.65 | -0.39 |
| w/ radiolarian test wall maximum $\delta^{30}\text{Si}$ | -1.50 | -0.82 | -1.1  | -0.98 |
| <b><math>\delta^{18}\text{O}</math> analyses</b>        |       |       |       |       |
| <i>radiolarian interior</i>                             |       |       |       |       |
| n analyses  |       | 10    | 2     |       |
| median (‰)  |       | 26.0  | 22.6  |       |
| standard deviation (‰)                                  |       | 1.1   | 2.7   |       |
| maximum (‰)   |       | 27.2  | 24.5  |       |
| minimum (‰)   |       | 24.2  | 20.6  |       |
| <i>radiolarian test wall</i>                            |       |       |       |       |
| n analyses  |       | 23    | 6     |       |
| median (‰)  |       | 26.1  | 26.4  |       |
| standard deviation (‰)                                  |       | 2.1   | 1.1   |       |
| maximum (‰)   |       | 30.4  | 26.8  |       |
| minimum (‰)   |       | 22.1  | 25.8  |       |
| <i>spicule</i>  |       |       |       |       |

|                        |  |       |      |  |
|------------------------|--|-------|------|--|
| n analyses             |  | 6     | 6    |  |
| median (‰)             |  | 28.2  | 25.1 |  |
| standard deviation (‰) |  | 2.8   | 1.0  |  |
| maximum (‰)            |  | 31.7  | 26.0 |  |
| minimum (‰)            |  | 24.8  | 23.6 |  |
| <i>matrix</i>          |  |       |      |  |
| n analyses             |  | 24    | 35   |  |
| median (‰)             |  | 25.1  | 24.0 |  |
| standard deviation (‰) |  | 2.0   | 1.0  |  |
| maximum (‰)            |  | 27..5 | 26.3 |  |
| minimum (‰)            |  | 20.6  | 22.2 |  |

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